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CORRECTION

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Correction: cDNA-AFLP analysis reveals differential gene expression in compatible interaction of wheat challenged with *Puccinia striiformis* f. sp. *tritici*

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Text

After the publication of this work [1], we became aware of the fact that the GenBank accessions listed in column 2 of Table 1 and Additional File 1 are incorrect. The EF# accession numbers should be prefixed with FF# and not with EF#. The corrected Table 1 is provided below. The corrected additional file can be accessed via the additional file contained in this article.

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Table 1 Transcript derived fragments (TDFs) from wheat leaves infected by *Puccinia striiformis* f. sp. *tritici* with homologies to genes in *P. graminis* f. sp. *tritici*

TDF	Accession No.	Size (bp)	<i>P. graminis</i> f. sp. <i>tritici</i> clones	E-value
PST_72-1-2b	FF339818	235	PGTG_02587 Glycine dehydrogenase	7e-15
PST_C40	FF339683	311	PGTG_04973 Fructose-1,6-bisphosphatase	1e-28
PST_C37 ^a	FF339680	294	PGTG_01121 UDP-glucuronic acid decarboxylase	3e-22
PST_C38	FF339681	265	PGTG_15605 ATP synthase subunit alpha	3e-15
PST_C87	FF339730	237	PGTG_04870 ATP synthase subunit beta	1e-15
PST_C59 ^a	FF339702	643	PGTG_06894 NADH-quinone oxidoreductase chain 3	3e-74
PST_315-3 ^a	FF339797	407	PGTG_16250 possible glycosyl transferase	1e-16
PST_84-3b ^a	FF339823	410	PGTG_13068 Conserved hypothetical protein	7e-25
PST_C81	FF339724	318	PGTG_08200 Vesicular-fusion protein SEC17	3e-14
PST_C16	FF339660	366	PGTG_14848 Conserved hypothetical protein	2e-09
PST_C88	FF339731	216	PGTG_14274 Plasma membrane proteolipid 3	2e-12
PST_68b-1	FF339813	259	PGTG_18059 NADH-quinone oxidoreductase	4e-06
PST_68b-3 ^a	FF339814	257	PGTG_18059 NADH-quinone oxidoreductase	7e-16
PST_C101 ^a	FF339744	586	PGTG_07295 Conserved hypothetical protein	1e-21
PST_315-4 ^a	FF339798	404	PGTG_10913 Predicted protein	1e-21
PST_C86 ^a	FF339729	436	PGTG_15782 Hypothetical protein	8e-32
PST_C83 ^a	FF339726	414	PGTG_02587 Glycine dehydrogenase	7e-33
PST_C73	FF339716	249	PGTG_13068 Conserved hypothetical protein	2e-24

^a These genes are confirmed to be of stripe rust fungus origin by sequencing.
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Additional file

Additional file 1: Transcript derived fragments (TDFs) from *Puccinia striiformis* f. sp. *tritici* infected wheat leaves with altered expression patterns and their closest matches in the GenBank database.

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